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<150> GB 9825303.2
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35 40 45	
Ala Gln Leu Pro Gly Leu Arg His Leu Asp Leu Arg Asn Asn Ser Leu	
50 55 60	
Val Ser Leu Thr Tyr Val Ser Phe Arg Asn Leu Thr His Leu Glu Ser	
65 70 75 80	
Leu His Leu Glu Asp Asn Ala Leu Lys Val Leu His Asn Ala Thr Leu	
85 90 95	
Ala Glu Leu Gln Ser Leu Pro His Val Arg Val Phe Leu Asp Asn Asn	
100 105 110	
Pro Trp Val Cys Asp Cys His Met Ala Asp Met Val Ala Trp Leu Lys	
115 120 125	
Glu Thr Glu Val Val Pro Gly Lys Ala Gly Leu Thr Cys Ala Phe Pro	
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Glu Lys Met Arg Asn Arg Ala Leu Leu Glu Leu Asn Ser Ser His Leu	
145 150 155 160	
Asp Cys Asp Pro Ile Leu Pro Pro Ser Leu Gln Thr Ser Tyr Val Phe	
165 170 175	
Leu Gly Ile Val Leu Ala Leu Ile Gly Ala Ile Phe Leu Leu Val Leu	
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Ala Cys Arg Asp His Met Glu Gly Tyr His Tyr Arg Tyr Glu Ile Asn
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